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<120> Novel compounds

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<212> DNA

<213> *Neisseria meningitidis*

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<211> 400

<212> PRT

<213> *Neisseria meningitidis*

<400> 2

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 Val Thr Lys Ser Asp Leu Glu Glu Glu Gln Phe Glu Thr Arg Ile Gly
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 Gln Ser Val Ala Gly Val Ser Glu Asn Gly Asn Val Leu Ala Arg Leu
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 210 215 220
 Arg Ile Ala Pro Glu Pro Ala Gln Thr Asp Lys Gln Asp Ser Lys Ser
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 Ile Asn Leu Ala Leu Thr His Tyr Asn Asp Lys Gln Asp Thr Asp Tyr
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<211> 2409

<212> DNA

<213> *Neisseria meningitidis*

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<210> 4

<211> 802

<212> PRT

<213> *Neisseria meningitidis*

<400> 4

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Val Glu Gly Gln Ser Gln Val Lys Val Arg Ala Glu Gly Gly Val Val
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 Tyr Gly Gly Arg Ala Ala Gly Gly Ser Leu Asn Ala Gly Leu Ser Val
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<210> 5

<211> 2016

<212> DNA

<213> *Neisseria meningitidis*

<400> 5

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<210> 6

<211> 671

<212> PRT

<213> *Neisseria meningitidis*

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Ile Thr Arg Leu Lys Lys Leu Arg Asp Arg Asn Ser Glu Tyr Trp Lys
      50              55              60
Glu Glu Thr Tyr His Ile Lys Ser Asn Asn Arg Val Tyr Pro Asn Ile
65              70              75              80
Pro Ala Leu Phe Pro Lys His Pro Phe Asp Pro Phe Glu Asn Ile Asn
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Thr Glu Lys Pro Ile Arg Gln Tyr Phe Lys Glu Cys Leu Asn Thr Gly
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Lys Tyr Ser Asp Asp Thr Cys Lys Ser Gln Gln Ser Ile Pro Thr Val
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Arg Ser Asp Ile Phe Ala Leu Asn Thr Lys Ile Lys Asn Ser His Ile
      165             170             175
Asn Ser Glu Ile Phe Ala Val Gly Asn Tyr Thr Lys Leu Met Tyr Ser
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Ala Gln His His Ser Ile Trp Ser Glu His Leu Tyr Ser Asn Ser Glu

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Val Tyr Leu Gln Pro Gln Ala Gln Leu Thr Tyr Leu Gly Val Asn Gly		
545	550	555
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565	570	575

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<210> 7

<211> 2076

<212> DNA

<213> *Neisseria meningitidis*

<400> 7

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<211> 691

<212> PRT

<213> *Neisseria meningitidis*

<400> 8

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Met Thr Leu Lys Ala Ser Lys Gln Ala Ser Gly Arg Val Asn Leu Leu
 1          5          10          15
Thr Leu Ser Val Leu Ser Leu Phe Cys Thr Pro Tyr Val Cys Gly Ser
 20          25          30
Asp Ala Tyr Asp Pro Val Lys Glu Ala Glu Ile Lys Asn Lys Phe Ile
 35          40          45
Leu Glu Ala Ala Glu Asp Arg Asn Ser His Val Trp Arg Gly Pro Cys
 50          55          60
Ser Ile Ser Phe Asp Cys Phe Gly Met Phe Arg Ala Gln Leu Gly Ser
 65          70          75          80
Asn Thr Arg Ser Thr Lys Ile Gly Asp Asp Ala Asp Phe Ser Phe Ser
 85          90          95
Asp Lys Pro Lys Pro Gly Thr Ser His Tyr Phe Ser Ser Gly Lys Thr
100          105          110
Asp Gln Asn Ser Ser Glu Tyr Gly Tyr Asp Glu Ile Asn Ile Gln Gly
115          120          125
Lys Asn Tyr Asn Ser Gly Ile Leu Ala Val Asp Asn Met Pro Val Val
130          135          140
Lys Lys Tyr Ile Thr Asp Thr Tyr Gly Asp Asn Leu Lys Asp Ala Val
145          150          155          160
Lys Lys Gln Leu Gln Asp Leu Tyr Lys Thr Arg Pro Glu Ala Trp Glu
165          170          175
Glu Asn Lys Lys Arg Thr Glu Glu Ala Tyr Ile Glu Gln Leu Gly Pro
180          185          190
Lys Phe Ser Ile Leu Lys Gln Lys Asn Pro Asp Leu Ile Asn Lys Leu
195          200          205
Val Glu Asp Ser Val Leu Thr Pro His Ser Asn Thr Ser Gln Thr Ser
210          215          220
Leu Asn Asn Ile Phe Asn Lys Lys Leu His Val Lys Ile Glu Asn Lys
225          230          235          240
Ser His Val Ala Gly Gln Val Leu Glu Leu Thr Lys Met Thr Leu Lys
245          250          255
Asp Ser Leu Trp Glu Pro Arg Arg His Ser Asp Ile His Thr Leu Glu

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				260					265					270			
Thr	Ser	Asp	Asn	Ala	Arg	Ile	Arg	Leu	Asn	Thr	Lys	Asp	Glu	Lys	Leu		
		275						280				285					
Thr	Val	His	Lys	Ala	Tyr	Gln	Gly	Gly	Ala	Asp	Phe	Leu	Phe	Gly	Tyr		
	290					295					300						
Asp	Val	Arg	Glu	Ser	Asp	Glu	Pro	Ala	Leu	Thr	Phe	Glu	Gln	Asn	Val		
305					310					315					320		
Ser	Gly	Lys	Ser	Gly	Val	Val	Leu	Glu	Arg	Arg	Pro	Glu	Asn	Leu	Lys		
			325						330					335			
Thr	Leu	Asp	Gly	Arg	Lys	Leu	Ile	Ala	Ala	Glu	Lys	Ala	Asp	Pro	Asn		
		340						345					350				
Ser	Phe	Ala	Phe	Lys	Gln	Asn	Tyr	Arg	Gln	Gly	Leu	Tyr	Glu	Leu	Leu		
	355						360					365					
Leu	Lys	Gln	Cys	Glu	Gly	Gly	Phe	Cys	Leu	Gly	Val	Gln	Arg	Leu	Ala		
370						375					380						
Ile	Pro	Glu	Ala	Glu	Ala	Val	Leu	Tyr	Ala	Gln	Gln	Ala	Tyr	Ala	Ala		
385					390					395					400		
Asn	Thr	Leu	Phe	Gly	Leu	Arg	Ala	Ala	Asp	Arg	Gly	Asp	Asp	Val	Tyr		
			405						410					415			
Ala	Ala	Asp	Pro	Ser	Arg	Gln	Lys	Leu	Trp	Leu	Arg	Phe	Ile	Gly	Gly		
		420						425					430				
Arg	Ser	His	Gln	Asn	Ile	Arg	Gly	Gly	Ala	Ala	Ala	Asp	Gly	Arg	Arg		
	435						440					445					
Lys	Gly	Val	Gln	Ile	Gly	Gly	Glu	Val	Phe	Val	Arg	Gln	Asn	Glu	Gly		
	450					455					460						
Ser	Arg	Leu	Ala	Ile	Gly	Val	Met	Gly	Gly	Arg	Ala	Gly	Gln	His	Ala		
465					470					475					480		
Ser	Val	Asn	Gly	Lys	Gly	Gly	Ala	Ala	Gly	Ser	Tyr	Leu	His	Gly	Tyr		
			485						490					495			
Gly	Gly	Gly	Val	Tyr	Ala	Ala	Trp	His	Gln	Leu	Arg	Asp	Lys	Gln	Thr		
		500						505					510				
Gly	Ala	Tyr	Leu	Asp	Gly	Trp	Leu	Gln	Tyr	Gln	Arg	Phe	Lys	His	Arg		
	515						520					525					
Ile	Asn	Asp	Glu	Asn	Arg	Ala	Glu	Arg	Tyr	Lys	Thr	Lys	Gly	Trp	Thr		
	530					535					540						
Ala	Ser	Val	Glu	Gly	Gly	Tyr	Asn	Ala	Leu	Val	Ala	Glu	Gly	Val	Val		
545					550					555					560		
Gly	Lys	Gly	Asn	Asn	Val	Arg	Phe	Tyr	Leu	Gln	Pro	Gln	Ala	Gln	Phe		
			565						570					575			
Thr	Tyr	Leu	Gly	Val	Asn	Gly	Gly	Phe	Thr	Asp	Ser	Glu	Gly	Thr	Ala		
		580						585									

Figure 1 consists of seven histograms arranged horizontally, each representing a different network size N . The histograms are labeled $N=10$, $N=20$, $N=30$, $N=40$, $N=50$, $N=60$, and $N=70$ from left to right. The x-axis for all histograms is the number of contacts per node, n , ranging from 0 to 10. The y-axis represents the frequency of nodes with a given number of contacts. For $N=10$, the distribution is broad, with a peak at $n=1$ and significant frequencies at $n=0$ and $n=2$. As N increases, the distribution becomes increasingly narrow and more concentrated around $n=1$. By $N=70$, the distribution is very sharp, with almost all nodes having exactly one contact.

Asp Gly Glu Lys Gln Thr Leu Ala Gly Arg Thr Ala Leu Glu Gly Arg
645 650 655
Phe Gly Ile Glu Ala Gly Trp Lys Gly His Met Ser Ala Arg Ile Gly
660 665 670
Tyr Gly Lys Arg Thr Asp Gly Asp Lys Glu Ala Ala Leu Ser Leu Lys
675 680 685
Trp Leu Phe
690

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